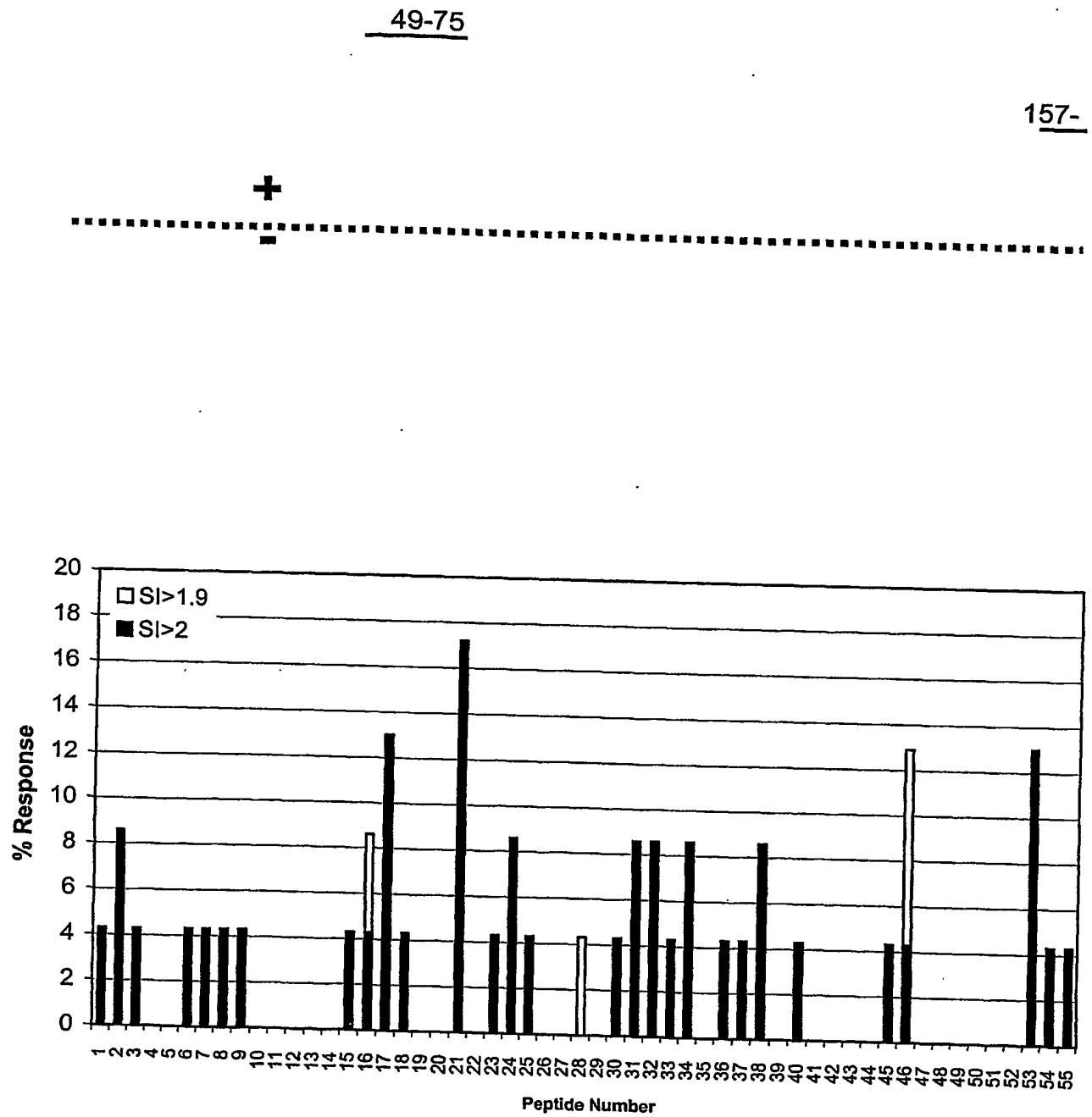


1 / 6

**Figure 1:****(A)****(B)**

SPAPPACDLRVL SKLLRDSHVLHS RLSQCPEVHPLP TPVLLPAVDFSL

2/6

1.

Region 1

GEWKTQMEETKA QDILGAVTLLE GVMAARGQLGPT CLSSLLGQLSGQ

49

VRLLLGALQSLL GTQLPPQGRRTA HKDPNAIFLSFQ HLLRGKVRFLML

97

Region 2

VGGSTLCVRRAP PTTAVPSRTSLV LTINEL

145

174

3 / 6

Figure 2A

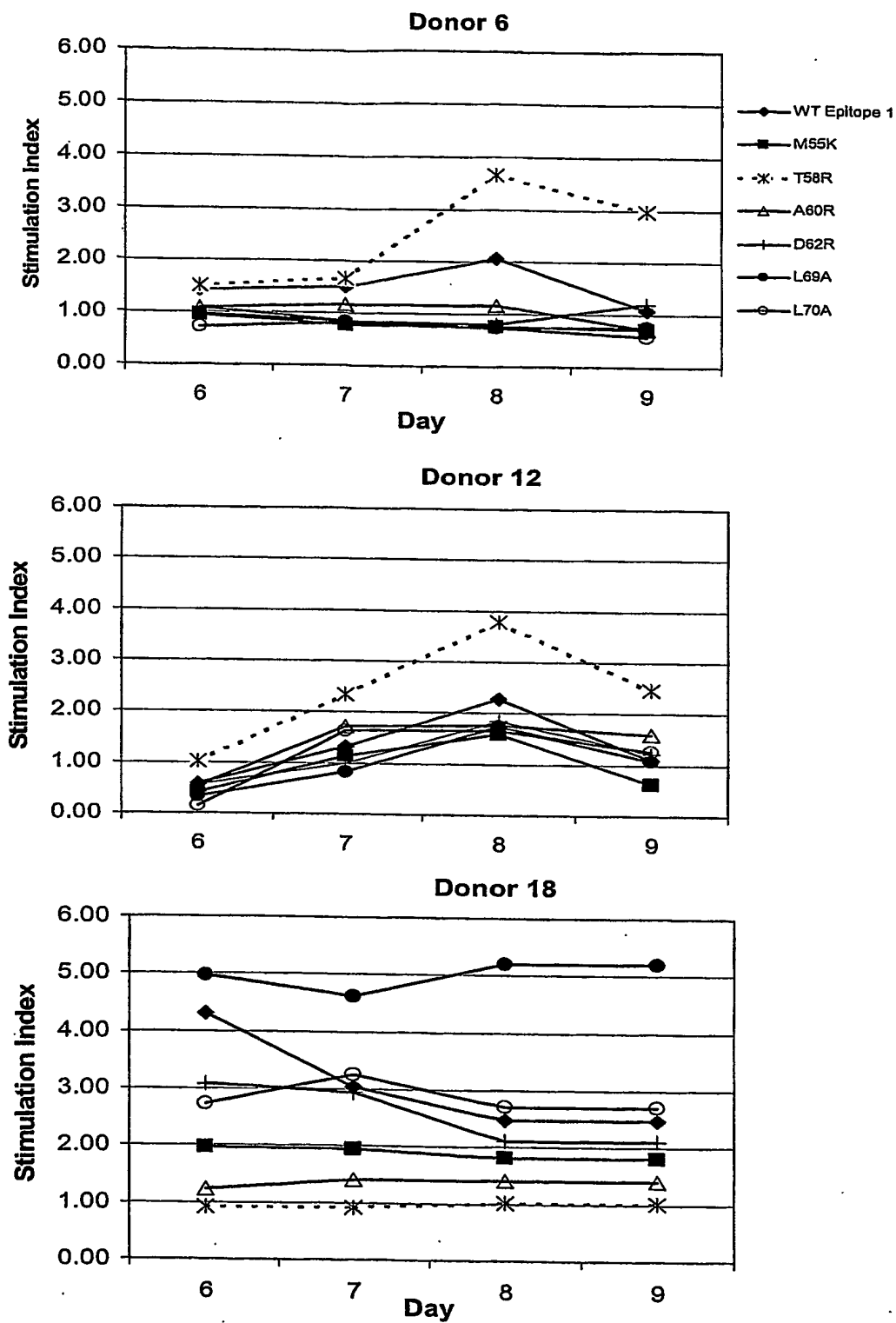
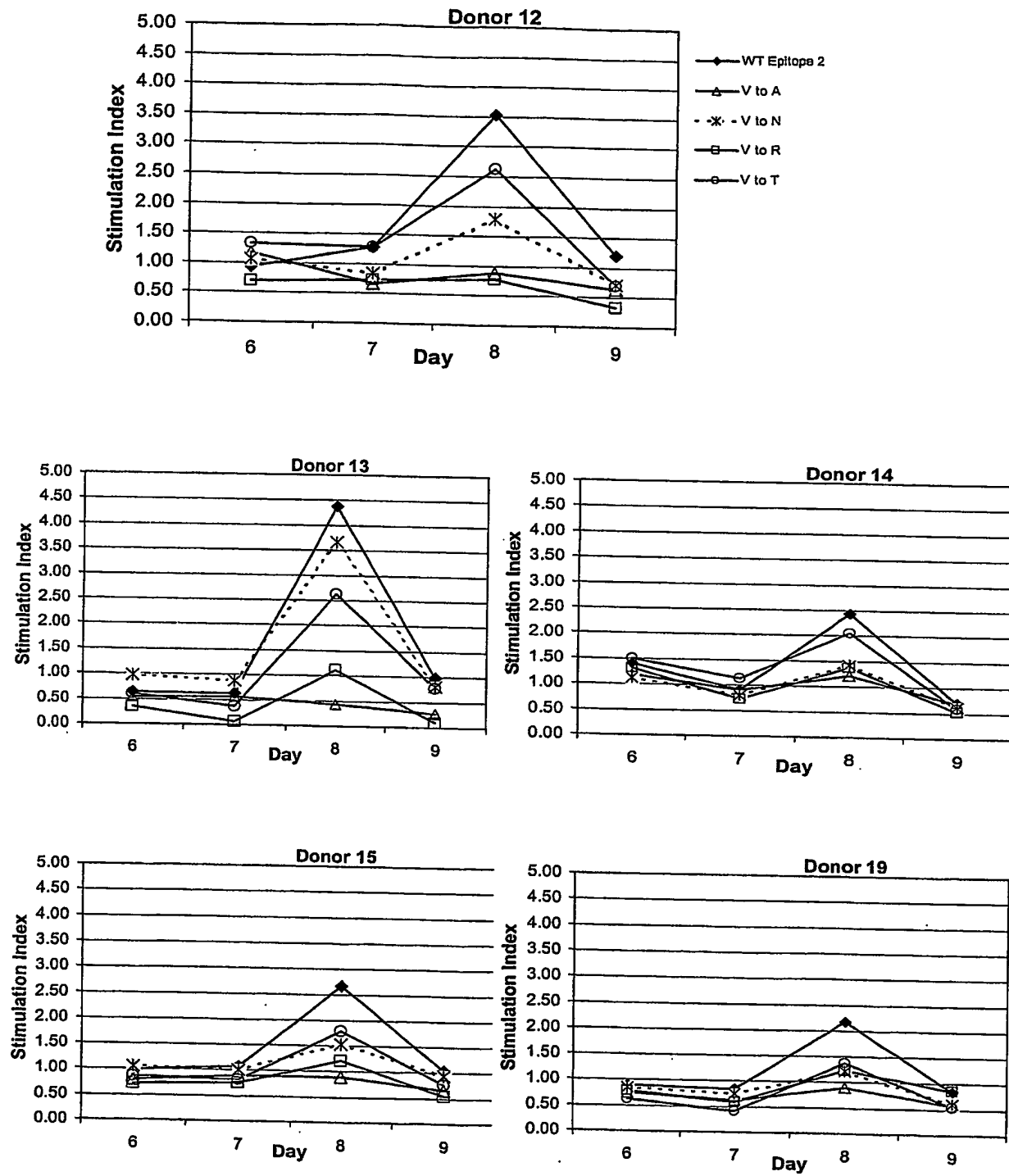
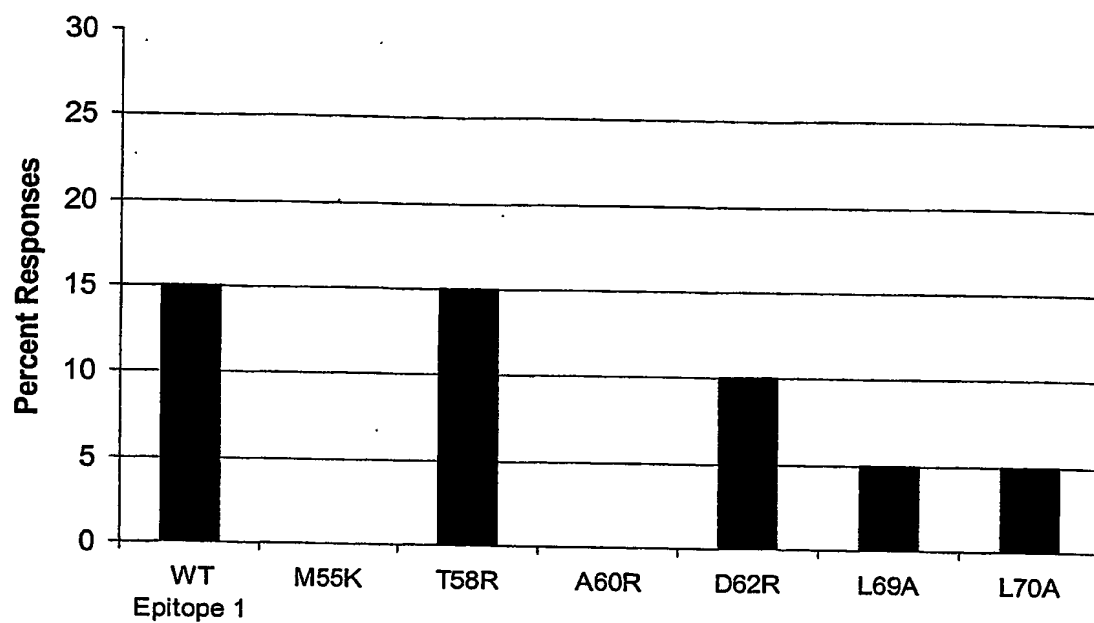
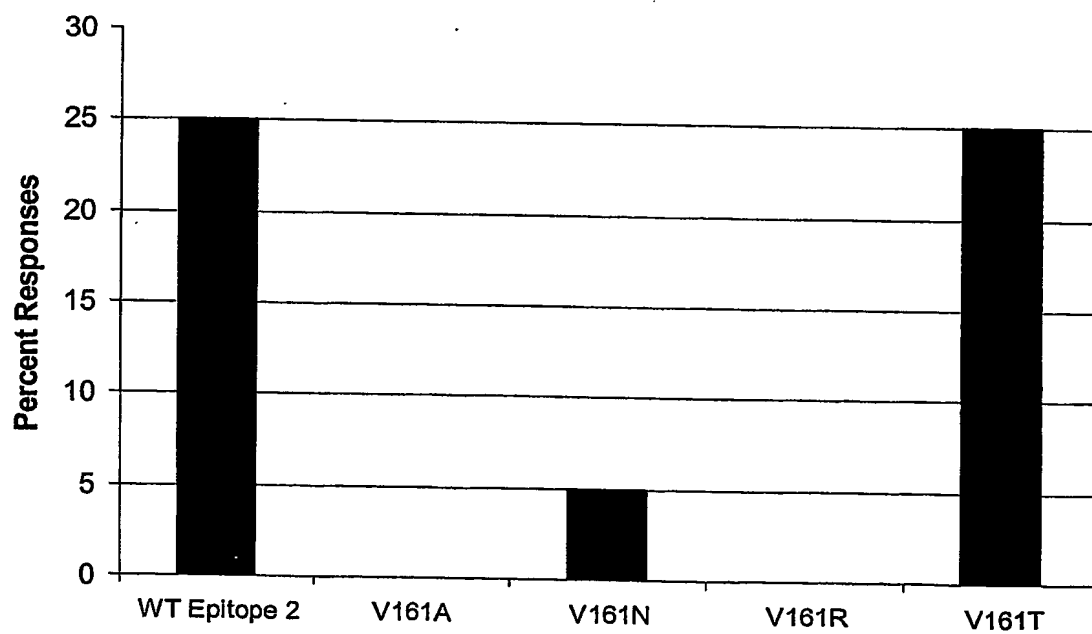


Figure 2B



**Figure 3****(A)****(B)**

**Figure 4**